## RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number:	10/572.175	
Source:	IFWP.	
Date Processed by STIC:	4/10/06	_
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## ENTERED

## **IFWP**

DATE: 04/10/2006 RAW SEQUENCE LISTING PATENT APPLICATION: US/10/572,175 TIME: 10:39:00 Input Set: A:\LeA 36 839.ST25.txt Output Set: N:\CRF4\04102006\J572175.raw 3 <110> APPLICANT: Bayer HealthCare AG Golz, Stefan Markova, Svetlana Burakova, Ludmila Frank, Ludmila Vysotski, Eugene 10 <120> TITLE OF INVENTION: Isolated Photoprotein mtClytin, and the Use Thereof 12 <130> FILE REFERENCE: LeA 36 839 C--> 14 <140> CURRENT APPLICATION NUMBER: US/10/572,175 C--> 14 <141> CURRENT FILING DATE: 2006-03-16 14 <150> PRIOR APPLICATION NUMBER: PCT/EP2004/009843 15 <151> PRIOR FILING DATE: 2004-09-03 17 <150> PRIOR APPLICATION NUMBER: DE10342670.1 18 <151> PRIOR FILING DATE: 2003-09-16 20 <160> NUMBER OF SEQ ID NOS: 9 22 <170> SOFTWARE: PatentIn version 3.3 24 <210> SEQ ID NO: 1 25 <211> LENGTH: 912 26 <212> TYPE: DNA 27 <213> ORGANISM: Clytia gregaria 29 <400> SEQUENCE: 1 60 30 gacagataaa aaattcactc cttagattat ttagtgaata agagaaaaaa aggataagaa 32 atcaagatgc aaaggtttac aaatcgtctt ctttccatgt cggctttacg tgcaagatca 120 34 agattgcaac gcacggcaaa ttttcacacc agcatactct tggctacaga ttcaaaatac 180 240 36 gcggtcaaac tcgatcctga'ttttgcaaat ccaaaatgga tcaacagaca caaatttatg 300 38 ttcaactttt tggacataaa cggtaagggg aaaatcacat tagatgaaat cgtctccaaa 360 40 gcttcagacg acatttgtgc taaactggat gcaacaccag aacagaccaa acgtcaccag 420 42 gatgctgttg aagccttttt caagaaaatg ggcatggatt atggtaaaga agttgcattc 480 44 ccagaattta ttaagggatg ggaagagttg gccgaacacg acttggaact ctggtctcaa 46 aacaaagta cattgatccg tgaatgggga gatgctgttt tcgacatttt cgacaaagac 540 48 gcaagtggct caatcagttt agacgaatgg aaggcttacg gacgaatctc tggaatctgt 600 660 50 ccatcagacg aagacgctga gaagacgttc aaacattgtg atttggacaa cagtggcaaa 52 cttgatgttg atgagatgac caggcaacat ttaggcttct ggtacacatt ggatccaact 720 54 tctgatggtc tttatggcaa ttttgttccc taagaagcgt tcagttaaaa acgctaaaca 780 56 ttgttcagtt gtaaaattat attcattttc atttcgtaaa attagtattt ataaatttgt 840 900 58 atcataaatt gtatccatgt tgtagactaa ataagactcg gcaaaaaaaa aaaaaaaaa 912 60 aaaaaaaaa aa 63 <210> SEQ ID NO: 2

64 <211> LENGTH: 228

68 <400> SEQUENCE: 2

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65 <212> TYPE: PRT

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79 35	40	-	45	
82 Pro Lys Trp Ile Ası	n Arg His Lys Phe	Met Phe Asn	Phe Leu Asp Ile	
83 50	55	60		
86 Asn Gly Lys Gly Lys	s Ile Thr Leu Asp	Glu Ile Val	Ser Lys Ala Ser	
87 65	70	75	80	
90 Asp Asp Ile Cys Ala	a Lys Leu Asp Ala	Thr Pro Glu	Gln Thr Lys Arg	
91 85		90	95	
94 His Gln Asp Ala Va	l Glu Ala Phe Phe	Lys Lys Met	Gly Met Asp Tyr	
95 100	105	5	110	
98 Gly Lys Glu Val Ala	a Phe Pro Glu Phe	e Ile Lys Gly	Trp Glu Glu Leu	
99 115	120	_	125	
102 Ala Glu His Asp Le	<del></del>	_		
103 130	135	140		
106 Arg Glu Trp Gly As	_			
107 145	150	155	160	
110 Gly Ser Ile Ser Le				
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114 Ile Cys Pro Ser As		_	<del>-</del>	
115 180	18 Ur Tra Ton Nan Va		190 The Ara Cla His	
118 Leu Asp Asn Ser G	ry Lys Leu Asp va 200	ar Asp Gru Me	205	
119 195 122 Leu Gly Phe Trp Ty		ro Thr Ser Ac		
122 heu Gry Flie 11p 1;	215	22)		
126 Asn Phe Val Pro	219	22	,	
127 225				
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132 <212> TYPE: PRT				
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162 aacggaaaaa tcactttgga tgaaattgtc tccaaagctt cggatgacat ttgcgccaaa
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164 cttggagcta caccagctca aacccaacgt catcaggaag ctgttgaagc tttcttcaag
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166 aagattggtt tggattatgg caaagaagtc gaattcccag ctttcgttaa cggatggaaa
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168 qaactqqcca aacatqactt qaaactttqq tcccaaaaca agaaatcttt gatccgcaat
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170 tggggagaag ctgtattcga cattttcgac aaggacggaa gtggctcaat cagtttggac
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172 gaatggaaaa catacggagg aatctctgga atctgtccat cagacgaaga cgctgaaaag
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174 accttcaaac attgcgattt ggacaacagt ggcaaacttg atgttgacga gatgaccaga
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176 caacatttgg gattctggta caccttggac cctaacgctg atggtcttta tggcaacttt
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186 <211> LENGTH: 198
187 <212> TYPE: PRT
188 <213 > ORGANISM: Clytia gregaria
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197
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200 Asp Ile Asn Gly Asn Gly Lys Ile Thr Leu Asp Glu Ile Val Ser Lys
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                                                   45
201
204 Ala Ser Asp Asp Ile Cys Ala Lys Leu Gly Ala Thr Pro Ala Gln Thr
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205
208 Gln Arg His Gln Glu Ala Val Glu Ala Phe Phe Lys Lys Ile Gly Leu
209 65
212 Asp Tyr Gly Lys Glu Val Glu Phe Pro Ala Phe Val Asn Gly Trp Lys
213
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                   85
216 Glu Leu Ala Lys His Asp Leu Lys Leu Trp Ser Gln Asn Lys Lys Ser
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217
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220 Leu Ile Arg Asn Trp Gly Glu Ala Val Phe Asp Ile Phe Asp Lys Asp
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221
224 Gly Ser Gly Ser Ile Ser Leu Asp Glu Trp Lys Thr Tyr Gly Gly Ile
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225
228 Ser Gly Ile Cys Pro Ser Asp Glu Asp Ala Glu Lys Thr Phe Lys His
229 145
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232 Cys Asp Leu Asp Asn Ser Gly Lys Leu Asp Val Asp Glu Met Thr Arg
233
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                                       170
                   165
236 Gln His Leu Gly Phe Trp Tyr Thr Leu Asp Pro Asn Ala Asp Gly Leu
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240 Tyr Gly Asn Phe Val Pro
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245 <211> LENGTH: 747
246 <212> TYPE: DNA
247 <213> ORGANISM: Clytia gregaria
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256 caaagetteg gatgacattt gegeeaaact tggageaaca ceagaacaga eeaaacgtea
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258 ccaggatgct gtcgaagctt tcttcaaaaa gattggtatg gattatggta aagaagtcga
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260 attcccagct tttgttgatg gatggaaaga actggccaat tatgacttga aactttggtc
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262 tcaaaacaag aaatctttga tccgcgactg gggagaagct gttttcgaca tttttgacaa
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264 agacggaagt ggctcaatca gtttggacga atggaaggct tatggacgaa tctctggaat
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266 ctgctcatca gacgaagacg ccgaaaagac cttcaaacat tgcgatttgg acaacagtgg
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268 caaacttgat gttgatgaga tgaccagaca acatttggga ttctggtaca ccttggaccc
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270 caacgctgat ggtctttacg gcaattttgt tccttaaaca tcgaaacaaa agcccaaaag
                                                                           660
                                                                           720
272 aagttttgga agaattattt gatactatca tttgttacta tttcgtaaca tgctatattt
274 tgtaacatgc tatatttaaa taatttt
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279 <212> TYPE: PRT
280 <213> ORGANISM: Clytia gregaria
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292 Asp Ile Asn Gly Asp Gly Lys Ile Thr Leu Asp Glu Ile Val Ser Lys
293
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296 Ala Ser Asp Asp Ile Cys Ala Lys Leu Gly Ala Thr Pro Glu Gln Thr
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297
300 Lys Arg His Gln Asp Ala Val Glu Ala Phe Phe Lys Lys Ile Gly Met
301 65
304 Asp Tyr Gly Lys Glu Val Glu Phe Pro Ala Phe Val Asp Gly Trp Lys
305
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308 Glu Leu Ala Asn Tyr Asp Leu Lys Leu Trp Ser Gln Asn Lys Lys Ser
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312 Leu Ile Arg Asp Trp Gly Glu Ala Val Phe Asp Ile Phe Asp Lys Asp
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316 Gly Ser Gly Ser Ile Ser Leu Asp Glu Trp Lys Ala Tyr Gly Arg Ile
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320 Ser Gly Ile Cys Ser Ser Asp Glu Asp Ala Glu Lys Thr Phe Lys His
321 145
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                        150
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324 Cys Asp Leu Asp Asn Ser Gly Lys Leu Asp Val Asp Glu Met Thr Arg
325
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328 Gln His Leu Gly Phe Trp Tyr Thr Leu Asp Pro Asn Ala Asp Gly Leu
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336 <210> SEQ ID NO: 9
337 <211> LENGTH: 198
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339 <213> ORGANISM: Clytia gregaria
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351	Asp	Ile	Asn	Gly	Asn	Gly	Lys	Ile	Thr	Leu	Asp	Glu		Val	Ser	Lys
352			35					40					45		_	
355	Ala	Ser	Asp	Asp	Ile	Cys		Lys	Leu	Gly	Ala		Pro	Ala	Gln	Thr
356		50					55					60			_	
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363	Asp	Tyr	Gly	Lys	Glu	Val	Glu	Phe	Pro	Ala	Phe	Val	Asn	Gly	_	Lys
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368				100					105			_	_	110		
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372			115		_			120	_			_	125			<b>-</b>
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376		130	٥				135	<u></u>		_		140			_ ^	
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384				_	165			_		170			_		175	
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VERIFICATION SUMMARY

DATE: 04/10/2006

PATENT APPLICATION: US/10/572,175

TIME: 10:39:01

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L:14 M:271 C: Current Filing Date differs, Replaced Current Filing Date